

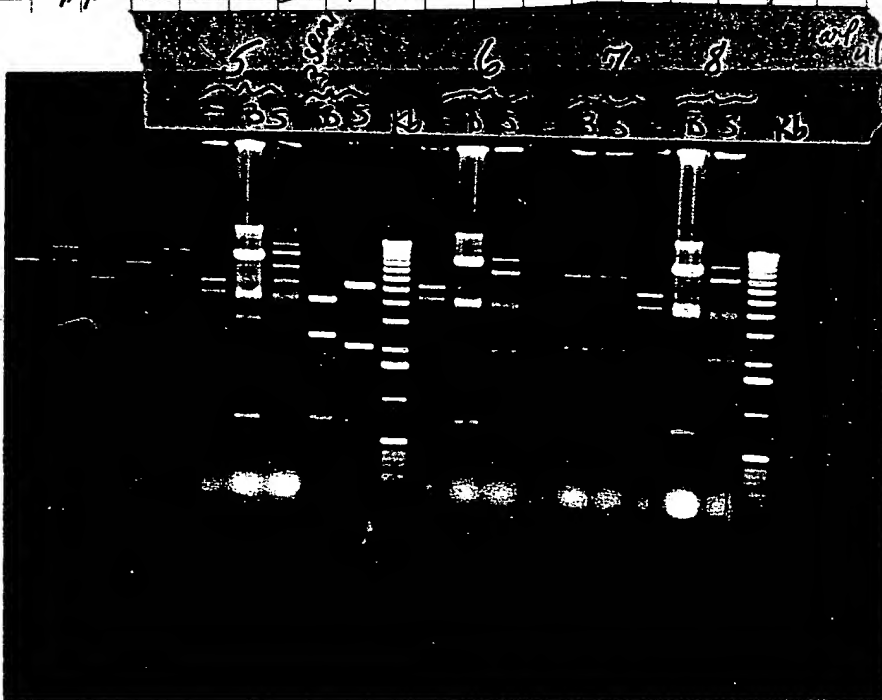
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Con'd from 3884 NB

2/18/95 wed

MINIPREP DNA

- cfr 500 μ l of cells for 1 minute in an eppendorf cfr (centrifuge)
- removed supernatant and resuspended pellet in 100 μ l of 1X PEBI (SI) ^(saved)
- added 200 μ l of alkaline - SDS mix
- placed the tubes on ice for few minutes (3-5 min)
- added 150 μ l of 7.5 M Ammonium Acetate
- mixed the tubes by inverting
- cfr the tubes for \sim 7-10 min
- transferred 400 μ l supernatant to the new eppendorf tube
- added 800 μ l of ethanol to supernatant. mixed tubes
- incubated the tubes for \sim 2 min. spin
- dissolved pellet in 50 μ l of TE + RNase A
- applied 5 μ l to a 1% agarose gel.



SI = 0.9% glucose

25 mM Tris HCl (pH 8.00)

10 mM EDTA

alkaline - SDS mix = 1% SDS

0.1 N NaOH

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2/9/95 The

Purification of m13 ssDNA

1. cfg 1.0 ml of *infected cell culture for 2 min. (1 to 5 min)
2. Transferred 800.0 μ l to the new tubes
(Pellet was saved for isolation of RF DNA)
3. cfg supernatant again to remove any residual cells
4. added 200.0 μ l of 20% PEG + 1.5 M NaCl. Vortexed
5. Incubated tubes at room temperature for 5 min.
6. cfg tubes for 5 min. & discarded supernatant (sup.)
7. added 200 μ l of *TE & vortexed really good.
8. cfg for ~ 1-2 min. (to remove any residual cell debris)
9. transferred sup. to the new tubes. (RNaseA can be added here)
10. added equal vol. of phenol / chloroform / isoamyl alcohol
(25:24:1) Mixed well.
11. cfg 5 min.
12. removed the aq. (upper) layer to a new tube (be very careful)
13. added $\frac{1}{10}$ vol. of 3M NaAc + $2\frac{1}{2}$ -3 vol. of 95% \downarrow
14. Incubated @ -70°C till 2/14/95.

 $\left\{ \begin{array}{l} 20.0 \mu\text{l NaCl} \\ 600.0 \mu\text{l Et} \end{array} \right.$ $\text{TE (T}_{10}\text{E}_1) = 10 \text{ mM Tris-HCl pH } 8.0 + 1 \text{ mM EDTA pH } 8.0$

infected cell culture = ① grew an E. coli F' strain to an OD of 0.4 in 2xYT
 \downarrow next pag
F' = Fertility Factor: codes for tra genes & pilis to allow infection of the
m13 Phage. (transfer of DNA)

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Cont'd

- ② Inoculated 1.0 ml of the cells ~~not~~ with the phage
- ③ Incubated the phage infected cells at 37°C for 5 hours. Now ^{was} supernatant ~~can be~~ processed for isolation of ssDNA & cells (pellet) were ready for isolation of RF (Replicating Form) dsDNA.

2/14/95 Tues.

1. Poured 0.8% Agarose gel (250 ml Volumes) in 1X TAE Buffer

2g Agarose

250.0 ml 1X TAE Buffer

- weight the flask
- boiled for 4.20 min. (brought up the)
- weight the flask & adjusted Volume to before boiling with • distilled water
- poured it on the plate.

2. 2X YT

added: 19.3g Tryptone } brought total Volume (TV)
 12g Yeast Extract } to 1200 ml with water
 12g NaCl }

- made 5 aliquots (1) 500.0 ml (2) 250 ml (3) 100.0 ml
 (4) 100.0 ml
 (5) 100.0 ml
- autoclaved at low pressure for 20 min.

3. 2X YT Top (soft) Agar

added: 0.35g Agar } made 3 different
 50.00 ml 2X YT } aliquots.

- autoclaved at low pressure for 20 min. (same as 2X YT)

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4. started 2, 10 ml cultures

#1 CJ236 in 2XYT + Cm 5 μ g/ml +/- phage from T. nea / m13
cloning at 37°C air-shaker

10 ml CJ236 in 2XYT

50 μ l Cm 5 μ g/ml 1000 μ g = mg/ml = 1000 μ l

#2 10 ml culture of T. nea / pTTG and T. nea / pTTG 118 in LB
+ 100 μ g/ml Ampicillin at 30°C air-shaker

added: 10 ml

5 μ l.

5. centrifuged (cf) ssDNA from DH5 α F'IQ (tubes left at -70°C, 2/9/95) for 10 min. at room temperature.

• discarded supernate

(a) Rinsed the precipitate (ppt.) with 70% EtOH. (removed any residual EtOH with another quick spin.

(b) Dried the DNA pellet at 55°C heat block

(c) Dissolved the DNA in 50.0 μ l of TE

(d) applied 5.0 μ l on a 0.8% Agarose gel (made today).
m13 ssDNA was used as a control

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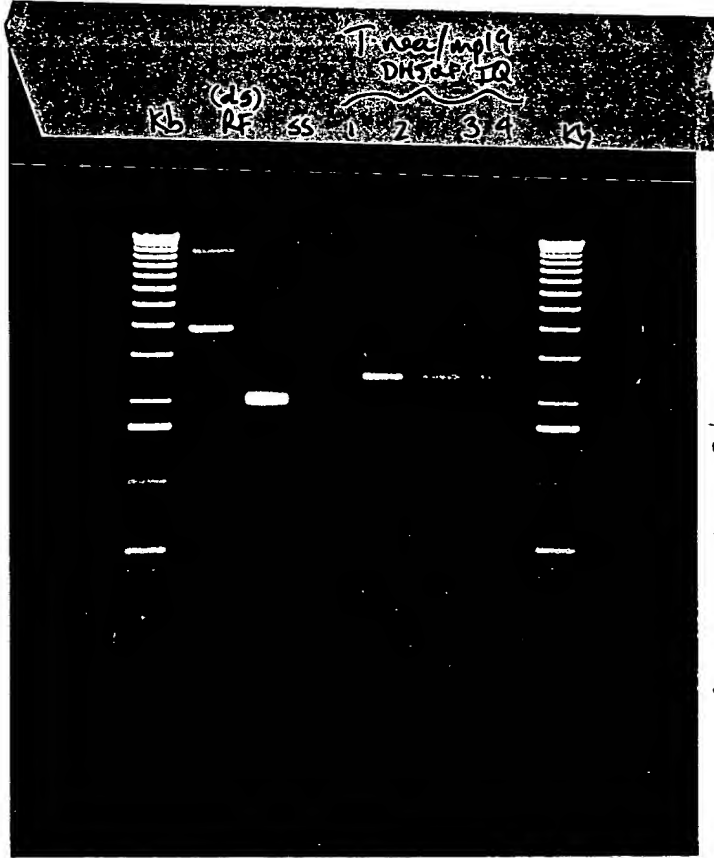


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RUN 140V ~ 2 hrs.

amp 4/12/95

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2/15/95 Wed.

(+) strand (ssDNA) lot # ED5702 260 μ g/ml
RF strand (dsDNA) lot # CC3111 5 μ g/18.4 μ l

calculation: ssDNA = 260 μ g/ml = ng/ μ l

$$\frac{260 \cancel{\mu\text{g/ml}}}{1000 \cancel{\text{ng}/\mu\text{g}}} \cdot 1000 \text{ ng}/\mu\text{g} \cdot \text{ml} \cdot 1000 \cancel{\mu\text{l}} = 0.260 \mu\text{g}/\mu\text{l}$$

$$\frac{1000 \text{ ng}/\mu\text{g} \cdot (0.260 \mu\text{g})}{260 \text{ ng}/\cancel{\mu\text{g}}} = 260 \text{ ng}/\mu\text{l}$$

$$\frac{260 \text{ ng}}{2.6} = 100 \text{ ng}$$

$$\left\{ \begin{array}{l} 260 \left(\frac{1}{2.6} \right) = 100 \text{ ng} \\ \text{or} \\ \frac{260}{2.6} = 100 \text{ ng} \end{array} \right.$$

for 2.6 total or final volume
you need 1.0 μ l DNA

$$\therefore \frac{1 \mu\text{l DNA} (260 \text{ ng}/\mu\text{l})}{1.6 \mu\text{l TE}} = 2.6 \mu\text{l}$$

for 100 ng/ μ l, } 2.0 μ l DNA (260 ng/ μ l)
multiply by 2 } 3.2 μ l TE

dsDNA = 5 μ g/18.4 μ l.

$$1000 \text{ ng}/\mu\text{g} \times 5 \mu\text{g} = \frac{1000 \text{ ng}(5 \mu\text{g})}{\cancel{\mu\text{g}}} = 5000 \text{ ng}/18.4 \mu\text{l}$$

$$\frac{5000 \text{ ng}}{18.4 \mu\text{l}} = \frac{272 \text{ ng}/\mu\text{l}}{2 \mu\text{l}} = 2.72 \text{ ng}$$

for 2.7 total volume you need
1.0 μ l DNA

Total Volume (TV)

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	RF (ds)	Tube # 1	Tube # 2
DNA	1.0 μ l $\times 3 = 3.0 \mu$ l		(+) ssDNA DNA = 1.0 μ l $\times 3 = 3.0 \mu$ l
TE	1.7 μ l $\times 3 = 5.1 \mu$ l		TE = 1.6 μ l $\times 3 = 4.8 \mu$ l
TV	2.7 μ l $\times 3 = 8.1 \mu$ l		TV = 2.6 μ l $\times 3 = 7.8 \mu$ l

Tube # 1, 2, 3, 4 of RF (dsDNA)

	① Alu I	② Hind III	③ Sau 3 A I	④ Bam HI	
H ₂ O	16.0 μ l				(all 4 tubes w/ 16.0 μ l)
10x Buffer	2.0 μ l				
DNA	1.0 μ l				
	(React 1; React 2; React 4; React 3)				
Alu I	+	-	-	-	
Hind III	-	+	-	-	
Sau 3 A I	-	-	+	-	
Bam HI	-	-	-	+	

Tube # 1, 2, 3, 4 of + (ssDNA) same order as RF

2 tubes were set-up for uncut, 1 with RF & 2nd with (+)

- each tube added 16.0 μ l H₂O
2.0 μ l React 2 10x buffer
1.0 μ l DNA

Put all 10 tubes in

- ran the sample (all 10) on a gel next morning.
(0.8% agarose gel, 147 volts)
- picture of the gel is on the next pg (pg # 8)

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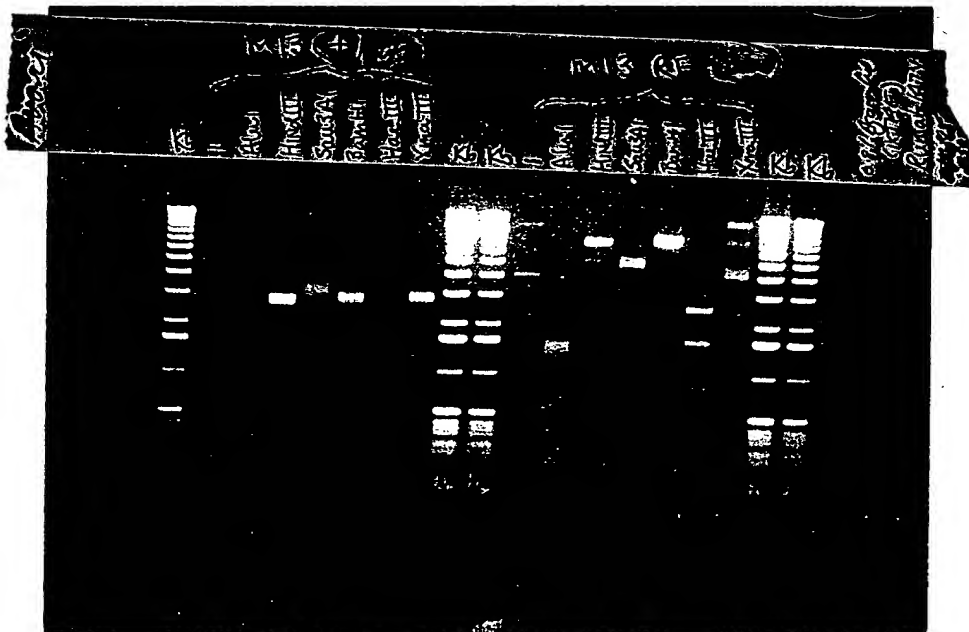
tube# 1 T. neu 1 pTTC

1.0 ml

tube# 2 T. neu 1 pTTC

1.0 ml

- Cfg. for 1 min. at room temperature
- discarded supernate and added: 100 μ l S1 to the pellet. mixed
200 μ l S2 lysis put both tube
ice.
150 μ l S2 with RNASE A
- Cfg. for 5 min. at 4°C
- transferred 400 μ l of supernatant to the new tubes.
- added 800.0 μ l E+OH to the supernatant
- put both tubes in the fridge till tomorrow (2/16/95)



arp 4/12/95

(+) Sal3A1 - gel shift (did not cut but binded)

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2/21/95 TUE

DIGEST T.nea/pSPORT with SstI & SphI

DIGEST M13 mp18 & M13 mp19 w/ SstI & SphI

M13 mp18 RF (0.44 ug/ul) } cut 500.0 ng
M13 mp19 RF (350.0 ug/ml)

$$\rightarrow 1000 \text{ ng/ug} \times 0.44 \text{ ug} = 440 \text{ ng}$$

$$\frac{500 \text{ ng}}{440 \text{ ng}} = 1.1 \text{ ul}$$

$$\rightarrow 1000 \times 0.350 \text{ ug/ul} = 350 \text{ ng}$$

$$\frac{350}{500 \text{ ng}} = 1.4 \text{ ul}$$

mp18

H₂O - 35.0 ul
10x buffer - 2.0 ul ← REact 2
500 ng DNA - 1.1 ul
1 ul SstI - 2.0 ul
40.0 ul

mp19

H₂O - 35.0 ul
10x buffer - 2.0 ul
DNA - 1.4 ul
SstI - 2.0 ul
40.0 ul

T.nea/pSPORT

H₂O - 81.0 ul
10x buffer - 10.0 ul
ng/ul DNA - 4.0 ul
SstI - 5.0 ul
100.0 ul

- Incubated all 3 tubes @ 37°C for 1/2 hour
- Made 0.8% agarose gel
250.0 ml TE buffer
2.0 g Agarose
- boiled for 4.0 min.
- added 12.0 ul E. Bio mide
- Poured the gel.

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added: 2.0 μ l of 1 M KCl 2 μ g = 2000 ng
to 40.0 μ g (mp 18 & 19)

5.0 μ l of 1M KCl
to 100.0 μ g (pSPORT)

added	Sph Sph I	-	2.0 uL	mp 18
			2.0 uL	mp 19
			5.0 uL	pSPORT

- Incubated @ 37°C for $\frac{1}{2}$ hour
- put the tubes in the fridge till
- ran samples on the gel ~~to see~~ on 2/22/95



arp 2/22/95 (1)

M13mp18 & mp19 RF D
are ds, supercoiled forms
the DNAs of phages M13
& 19. Using this vector
foreign DNA can be
inserted into the mul
cloning site in an
oriented fashion.

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2/22/95

1. grow cells overnight (O/N) 10.0 mL

= 9.0 mL (1.0 mL in ea. nine tubes). Each tubes labelled DH10B

• Quick freeze all nine tubes in a powdered
dry ice.

Plz C/T. nea

2/22/95 BJS

LB + AP100

GENE CLEAN

2) Did electrophoresis of yesterday's DNA (2/21/95)

M13 mp 18 and M13 mp 19 and pSPORT

b) Took the picture of the gel

c) cut off mp 18 fragment, mp 19 fragment & pSPORT fragment from
the gel & transformed the gel w/ ^{pa} DNA into the separate
Eppendorf tubes.d) added 700.0 μ L NaI to each ² tubes. Vortexed mp 18 & mp 19 tubes.e) Incubated both tubes @ 55°C to melt agarose. mixed ~~after~~ after incubation.f) added 5.0 μ L glass milk to both tubes.

g) Incubated both tubes on ice for 5 min.

h) Cfg. both tubes (quick spin)

i) discarded supernate

j) added 500.0 μ L New Wash bufferk) discarded supernate & again added 500.0 μ L New Wash buffer.
washed both tubes 3 times.l) added 10.0 μ L dH₂O to the tubes. mixed well by vortexing. 55°C for
2-5 min

m) set up Ligation

Ligation

H₂O = 12.0 μ Lligase) 5x Buffer = 4.0 μ Lmp 18 DNA = 2.0 μ L(1 μ /4L) Ligase = 2.0 μ LTV = 20.0 μ LH₂O = 12.0 μ L5x buffer = 4.0 μ Lmp 19 DNA = 2.0 μ LLigase = 2.0 μ LTV = 20.0 μ L

n) Incubated both tubes overnight @ room temperature (cond)

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(cont'd)

T.nea/Ptarc E1 pttc

- 1.0 mL of ea.
- Cfg.
- discarded supernate
- added 100.0 μ l S1 mixed well
- Incubated on ice for few min.
- added 200.0 μ l S2 14SIS
- Incubated on ice for few min.
- added 150.0 μ l S3 w/ RNAase A
- Cfg. for 7.0 min. @ 4°C

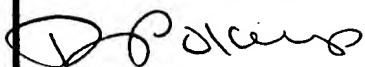
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2/28/95 TUE

I. set up digest DNA ppt.

① M13mp 18, ② mp 19 and ③ T. nea/pSPORT

1. - To ea. 3 added 100.0 μ L TE } to ppt.
" " 10.00 μ L NaAc } DNA
" " 300.00 μ L EtOH }

2. Incubated on dry ice for ~5 min.

3. Cfg. for 10 min @ room temp. (no ppt.)

4. no ppt., added 2.0 μ L (carrier molecule) Yeast tRNA. Vortexed

5. incubated on dry ice for ~5 min.

6. Cfg for 10 min. @ room temp. (Supernate saved) Pellet was saved on mp 18

7. added 200.0 μ L 70% EtOH to the pellet

8. Cfg. discarded supernate, air dried by putting tubes on the heat block.

II. DIGEST set-up. (to map Bam HI site)- cut T. nea/pSPORT with Hind III, Bam HI, Xba, NOT I, Sst, Eco R
SeparateH₂O - 13.0 μ Lbuffer - 2.0 μ LT. nea/pSPORT DNA - 3.0 μ Lenzyme - 2.0 μ LTV = 20.0 μ LEnzymes - Hind III, Xba, Sst had REact 2 1
buffer.- Bam HI, NOT I, Eco RI had REact:
- buffer.Control: H₂O - 13.0 μ L(REACT 2) buffer - 2.0 μ LDNA - 3.0 μ Lfor
separate
enzymes.

- Incubated @ 37°C

- ran on the gel on 3/1/95 (Wed)

Picture shown pg 21

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3/12/95

① 1 kb ladder ② T-neal/pSPORT uncut, ③ Sst, ④ Sst/sph, ⑤ sph, ⑥ 1 kb ladder (from 263 added loading dye, electrophoresis @ 190 v

- ~~digested~~ double digested BamHI/sphI (to map the Bam site T-neal

H₂O - 14.0 μ l

(REACT) buffer - 2.0 μ l

(T-neal/pSPORT) DNA - 2.0 μ l

(Bam/sph) enzyme - 1.0 μ l ea.

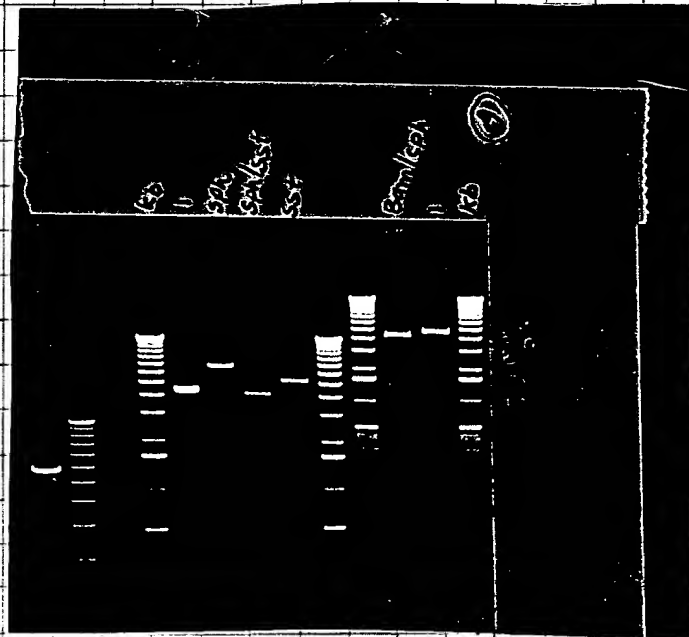
TV = 20.0 μ l.

control: - H₂O - 14.0 μ l

(uncut) buffer - 2.0 μ l

DNA - 2.0 μ l

Incubated @ 37°C for 30.0 min. (15 min.)



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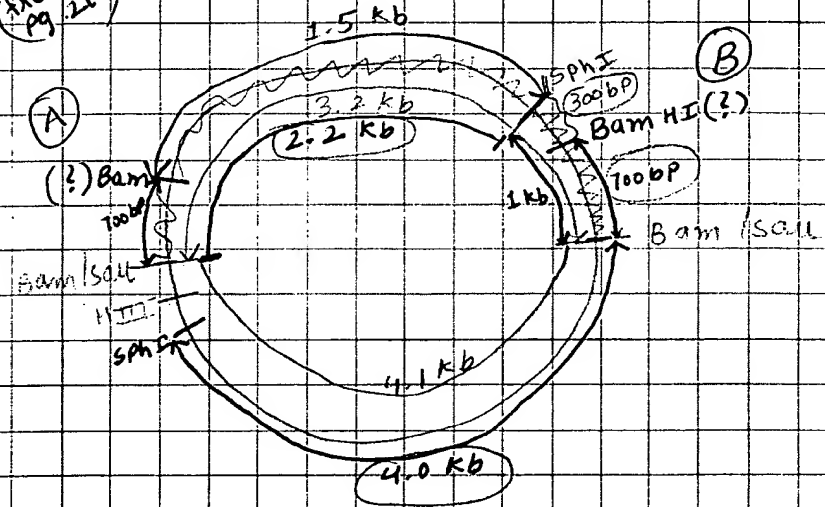
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Tag N. (from pg. 21)

GRAPH 2:



Bam I/Sph

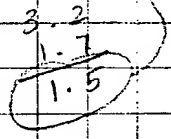
(A)

- 4.0 kb
- 700 bp
- 1 kb
- 1.5 kb

Bam I/Sph I

(B)

- 4.0 kb
- 2.2 kb
- 300 bp
- 700 bp

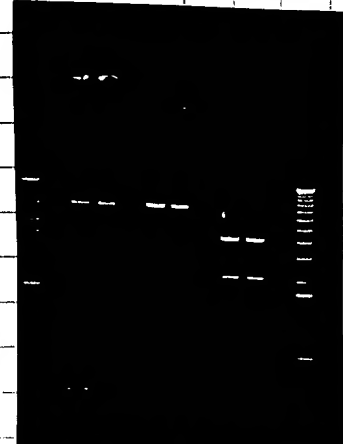


Cloning mp 18 w/ T. nea / pSPORT & mp 19 w/ T. nea / pSPORT

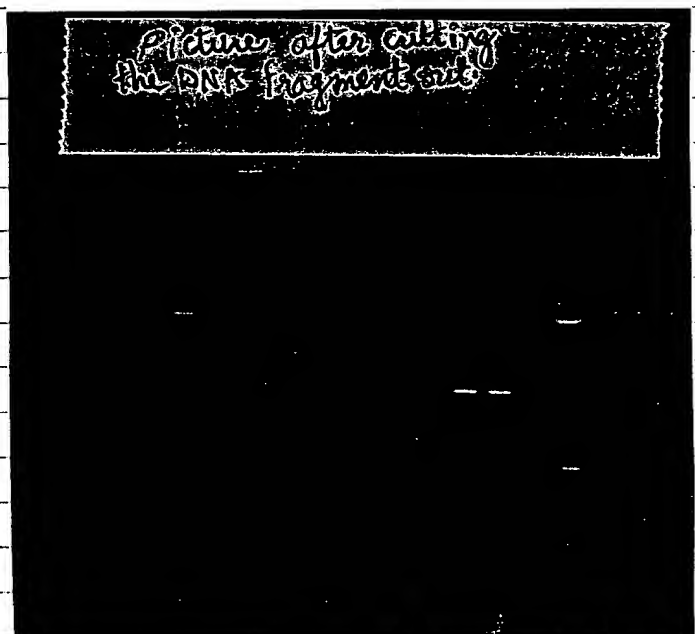
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~~3/2/95~~

Thurs.



Picture before cutting the DNA fragment.



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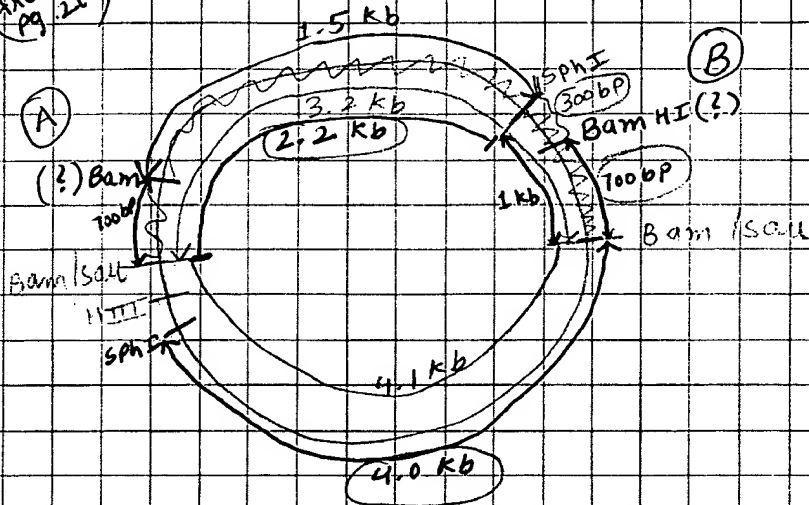
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ag N. — (SHOW)
pg. 21

GRAPH 2:



Bam 6 ph

(A)
4.0 Kb
700 bp
1 Kb
1.5 Kb

3.2
1.7
1.5

Bam SpH

(B)

4.0 kb

2.2 kb

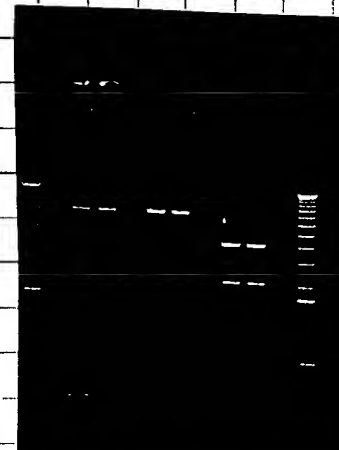
300 bp

700 bp

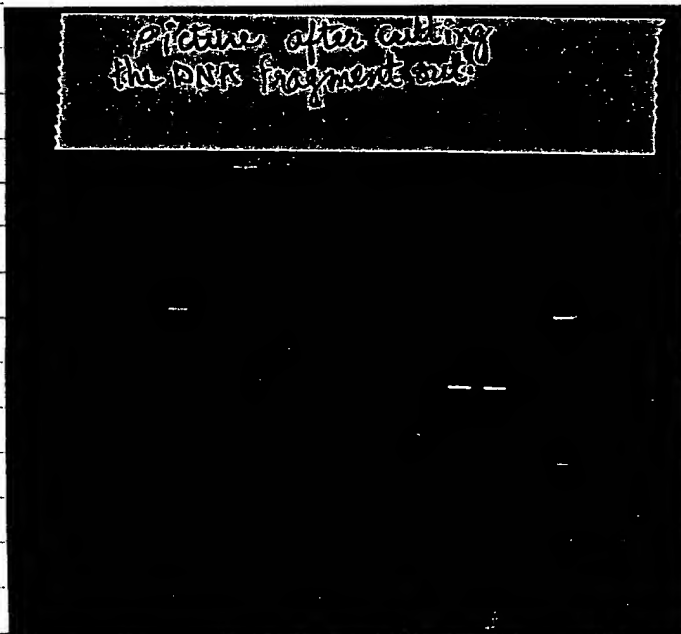
Cloning mp 18 w/ T.nea / pSPORT & mp 19 w/
T.nea / pSPORT

 $3 \mid 2 \mid 95$ ~~312195~~

Thurs.



Picture before
cutting the
~~DNA~~ DNA
fragment.



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GENE CLEAN

- (out) mixed mp 18 with T.nea pSPORT cut w/ Sst I/sph } 1 tube
- " mp 19 with " " " " } 2 tube

- added 700.0 μ l NaI to each 2 tubes. Vortexed
- put the tubes in 55°C heat block to melt agarose
- after agarose melted, added 5.0 μ l glass milk to both tubes
- incubated both tubes on ice for 5.0 min.
- c.f.g. both tubes (quick spin)
- discarded supernate & washed pellet 3 x with New Wash b
- added 14.0 μ l d H_2O to each tube
- quick spinned, discarded pellet & saved supernate

Set-up Ligation

- (mp 18) (mp 19) DNA - 14.0 μ l
- (ligase) 5x buffer - 4.0 μ l
- ligation - 2.0 μ l.
- TV - 20.0 μ l.

- incubated both

xfection cells

- (1) 100.0 μ l (competent)
- 3.0 μ l DNA (from ligation)

- (2) incubated on ice for 30 min.

- (3) heat shocked @ 42°C H_2O bath for 35 sec.

- melted 0.7% 2x YT top agar, added 4.0 μ l to 6 different glass tubes & put the tubes @ 55°C heat block

Didn't work

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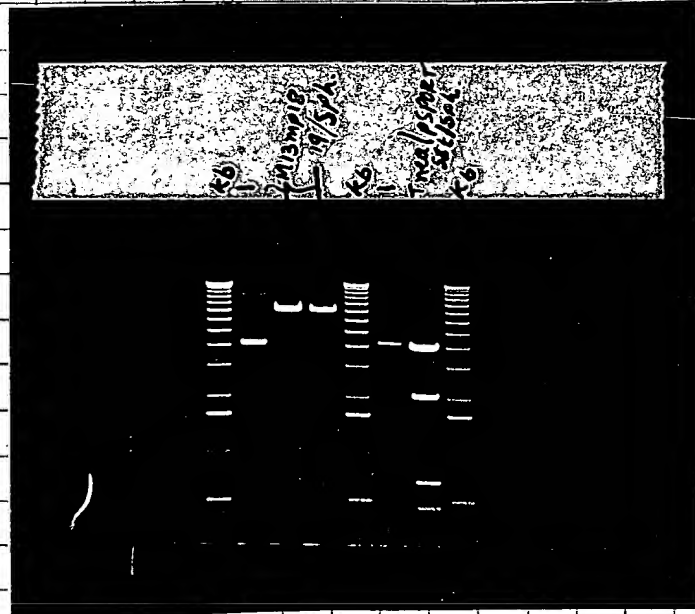
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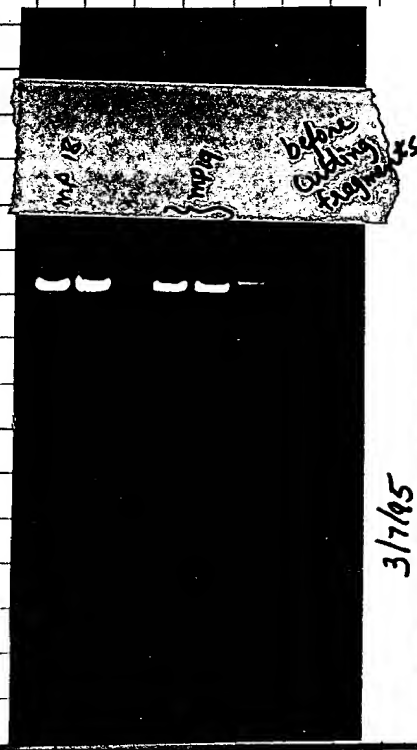
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 Repeat

3/7/95 TUE



3/7/95

After taking picture on looking @ the gel, ~~M±~~ M±3 mp18 and M±3 mp19 is @ the 7.2 Kb ~~cut~~ which was cut with Sph I. ~~we decided to cut~~ We planned on cutting mp18 and mp19 with Sst I. The gel ^{picture} below shows mp18 & mp19 before & after cutting the DNA fragments. After cutting the fragments performed Gene CLEAN



3/7/95



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		Recorded by <i>Duan</i>	

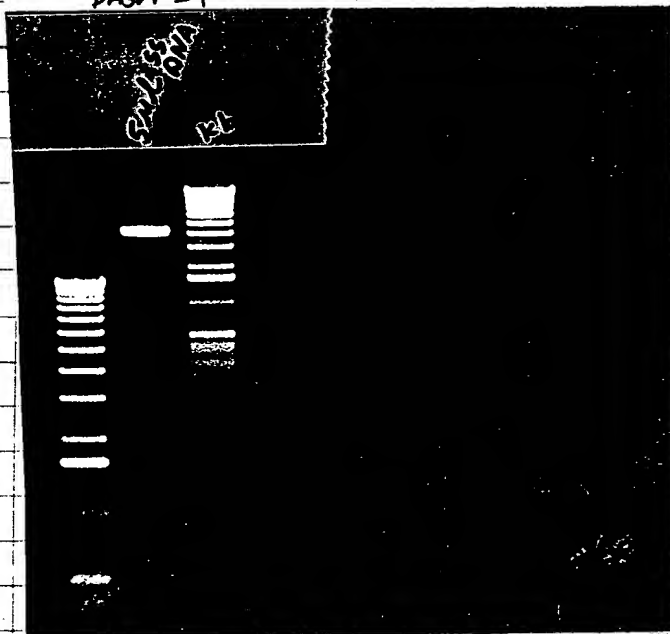
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labelled 2 tubes, 1 w/ mp 18, & 2nd w/ mp 19

1. to the DNA w/ agarose gel, added 700.0 μ l NaI
 2. put the tubes @ 52°C heat block to melt agarose. vortexed constant
 3. added 5.0 μ l glass milk to both tubes - mixed
 4. incubated on ice for 5 min.
 5. (fg. (quick spin) @ room temp.
 6. discarded supernate, added 500.0 μ l New wash buffer
 7. discarded supernate, washed pellet 3X with New wash buffer
 8. after washing 3X, added 14.0 μ l dH₂O to the pellet (discarded supernate (mixed))
 9. incubated @ 52°C for 5 min.
 10. discarded pellet & saved supernate for ligation.
- (could this on 3/8/95 wed.)

Purification of m13 ssDNA (T-neo 2kb [Sph1] / mp19) from pg. 1

T-neo/mp19 ssDNA (Sph1)
DH5 α 50



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ag N .

cell growth & Infection

- Grew an E. coli F' strain to an OD of 0.2-0.4 in 2x YT
- Inoculated 1-2 ml of the cells w/ the phage. (added 10.0 μ l from a liquid phage stock & added to cells)
- Incubated the phage infected cells @ 37°C for 5-7 hours.
- The supernate can now be processed for isolation of ssDNA & the cells can be processed for the isolation of Replication Form (RF) dsDNA.

Purification of ml3 ssDNA

- transferred 1.0 ml culture of infected cell to 4 different eppendorf tubes
- cfg 4 tubes for 2 min.
- transferred supernate to the new tubes & saved pellet from 1 tube (out of 4 tubes) for isolation of RF DNA
- Spinned the supernate again & transferred the supernate to the new tubes (done to remove any residual cells remained behind)
- passed the supernate through a 0.45 μ filter as to remaining cells (done when performing site-directed mutagenesis)
- added 200.0 μ l of 20% PEG + 1.5 M NaCl. Vortexed
- Incubated tubes for 15 min @ room temperature (or overnight @ 4°C)
- cfg. for 10 min in a μ cfg. @ room temp.
- discarded supernate & briefly spinned the tubes to remove the residual soln from the side of the tube (removed as much ^{supernate} as possible)
- added 200.0 μ l TE. Vortexed
- cfg for 2 min. to remove any residual cell debris.
- Transferred supernate to the new tube. (added 5.0 μ l RNase I to remove any residual nucleic acid from the prep. Benzonase will remove both RNA & DNA very efficiently.)
- added equal volume of phenol / chloroform / isoamyl alcohol mixed well
- cfg for 5.0 min.
- transferred the upper layer to a new tube (BE CAREFUL NOT TO DISTURB WHITE INTERFACE OR REMOVE ANY PHENOL)
- added 20.0 μ l NaAC & 600.0 μ l EtOH
- Incubated @ -70°C for 5-15 min. (we left @ -70°C overnight)

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3/8/95 wed

- cftg the samples for 10-15 min.
- discarded the supernate & rinsed the pellet w/ 70% EtOH
- dried the pellet @ 55°C heat block or @ room temperature
- dissolved the DNA in 50.0 μ l TE.

Shm Rxn

3/8/95 wed

Annealing Rxn.

+ Primer (2899)

- Primer (2899)

H₂O - 3.0 μ l4.0 μ l5x Buffer 2.0 μ l2.0 μ l19.5% SS DNA 4.0 μ l4.0 μ l(200mg/4L Kinasol) Oligo 1.0 μ lTV 10.0 μ l10.0 μ l

Incubated @ 70°C - 75°C for 2 min. (to eliminate non-spf. bin)
" @ 37°C - 40°C for 2 min.

Synthesis RxnAnnealing Rxn - 10.0 μ l.5mL 10x buffer - 2.0 μ lH₂O - 6.0 μ lT₄/T₇ DNA pol - 1.0 μ lT₄ DNA ligation - 1.0 μ lTV - 20.0 μ l.

Incubated @ 37°C for 10 min.

Synthesis rxn - 2.5 μ lTE - 8.0 μ lloading dye - 1.0 μ l

- ran the sample on the gel
- the picture on the next page, # 29.

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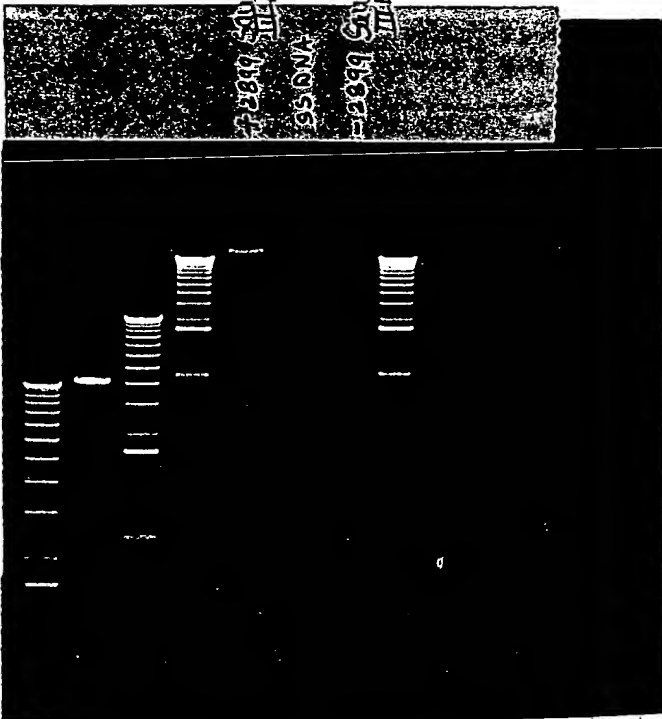
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4/12/95

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1289 5' 3' 1289 5' 3' 1289 5' 3'



(con'd from pg. 28)

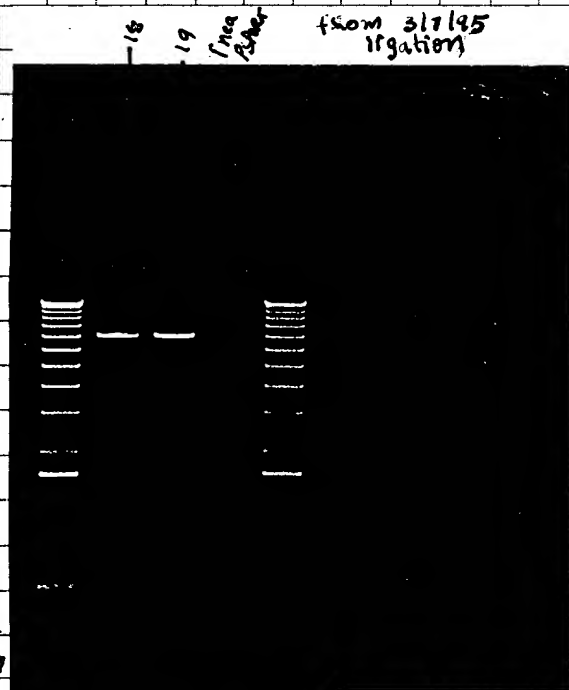
+2899 (w/ primer) oligo forms a ds DNA.
+2899 fragment looks brighter because Et. Bromide binds to it better. -2899 primer binds but ^(GWA) does not hold strongly.
∴ the DNA fragment looks fainter or light, less Et. Bromide is able to bind.

(con'd on pg. 41)

Ligation from 3/1/95 (pg. 26)

H₂O - 8.0 μ l
5X buffer - 4.0 μ l
mp 18 - 2.0 μ l
insert - 4.0 μ l
ligation - 2.0 μ l
TV - 20.0 μ l

H₂O - 8.0 μ l
5X buffer - 4.0 μ l
(vector) mp 19 - 2.0 μ l
insert - 4.0 μ l
ligation - 2.0 μ l
TV - 20.0 μ l



18, 19, ligation
3/1/95

- Incubated both samples for 1 hour @ room temp.

100.0 μ l Competent cells }
3.0 μ l DNA } xfection cells.

xfection

10% mp 18 / mp 19
90% mp 18 / mp 19
Control

ran mp 18 on 3/10
(used DNA from 3/10/95 again on 3/15/95 pg. 32)

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— 10% mp 18 / mp 19

added → 4.0 mL 2x YT TOP Agar

100.0 μ L X-Gal 4%5.0 μ L IPTG 200 mM (inducer = repressor gives tighter affinity)60.0 μ L lawn cells10.0 μ L x fraction cells (after heat shock for 35 sec.)

— 90% mp 18 & mp 19.

Same way as 10%

— Control

100.0 μ L X-Gal5.0 μ L IPTG60.0 μ L lawn cells.

no 3:30 to 4:00

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3/14/95 TUE

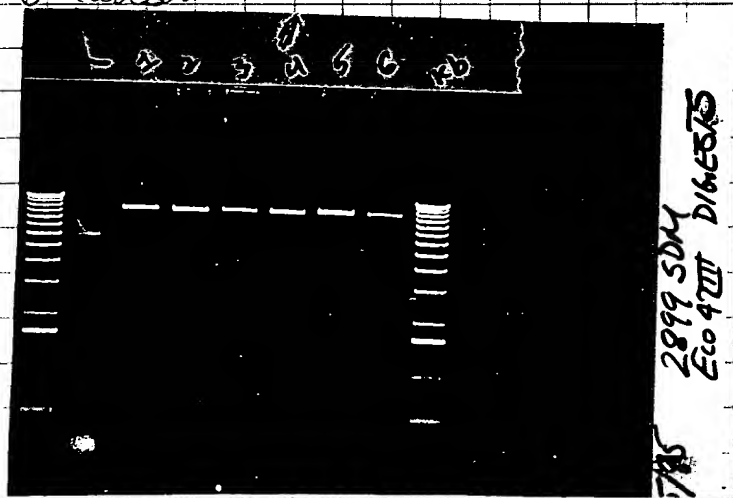
~~Reenza~~: miniprep

1.0 ml culture of T.nea /mp 19 grown for 5 hours @ 37°C in
 6 different glass tubes
 transferred 1 ml cell to the 6 different labelled eppendorf tubes.
 cfg all 6 tubes for 2 min. @ room temp.
 removed supernate & saved in different tubes
 added 100 μ l S1 mixed well
 added 200 μ l S2. put the tubes on ice (mixed by inverting)
 added 150 μ l 7.5 M NH_4OAc
 incubated on ice for 5 min.
 cfg for 7 min. @ room temp (4°C) NOTE: cfg in 4°C room was taken away for repair: used @ RT.
 transferred supernate (400.0 μ l) to the new 6 labelled tubes
 added 800 μ l of EtOH to the 400 μ l of supernate (mixed well)
 incubated @ -70°C for 30 min.
 cfg for 2 min. @ room temp (discarded supernate)
 rinsed w/ 70% EtOH (removed supernate)
 added 50.0 μ l TE to the pellet.

H_2O - 7.0 μ l	x 6	= 42.0 μ l
buffer - 2.0 μ l	x 6	= 12.0 μ l
Eco47III - 6.0 μ l	x 6	= 6.0 μ l
TV		60.0 μ l

added 10.0 μ l DNA⁺ to each 6 tubes.

the map is on next page # 32. Fragments
 on all 6 tubes are in still present,
 y haven't gone into the mutant.
 I tried miniprep again next day.
 (started)



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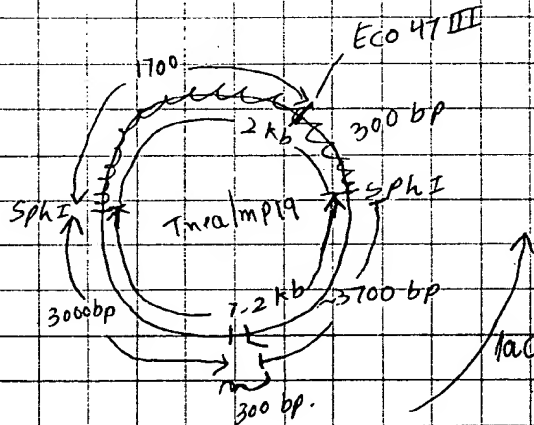
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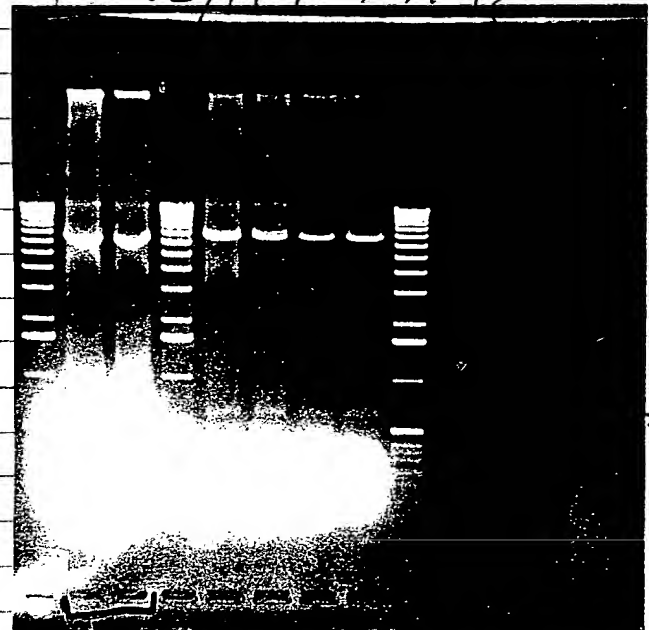
3/15/95 Wed



parent Eco 47 III
 ~ 8.9 kb
 0.3 kb → (most probably won't see fragment because too small & too light)

mutant
 4 kb
 4.7 kb
 0.3 kb

DNA from date 3/10/95



H₂O = 6.0 mL.
 R6 buffer = 2.0 mL.
 mpi8 DNA = 10.0 mL
 Sst/Sph 1.0 mL ea.
 TV 20.0 mL

H₂O = 6.0 mL.
 buffer = 2.0 mL
 mpi8 DNA = 10.0 mL
 Sst/Sph = 1.0 mL ea.
 TV 20.0 mL

from pg 29
 & ran again
 on 3/15

arp
 4/12/95

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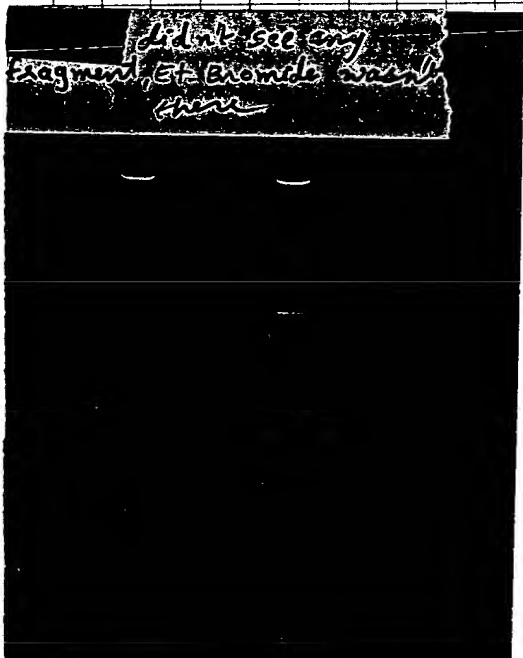
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- Incubated both tubes @ 37°C for 30 min.
- added 2.0 μL loading dye to each tube
- ran both on a gel
- took picture



3/15/95 T. nea/mp

1.0 ml T. nea (Sph I) / mp 19 + 2899 + Sau 3AI grown for 5 hours @ 37°C in 10 different glass tubes
 after 5 hours transferred 1.0 ml culture to the 10 labelled eppendorf tubes
 cfd all 10 eppendorf tubes @ room temperature for 2 min.
 removed supernate & saved
 put all 10 tubes w/ pellet & all 10 tubes w/ supernate @ -70°C overnight or until 3/16/95 Thursday.

TE: Brian had to leave @ 4:30 pm & this was a point to stop @.

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3/16/95 Thurs.

con'd from page 33 3/15/95 wed. MINIPREP

- took the pellet out from -70°C (10 eppendorf tubes)
- added 100 μL S1 mixed well
- added 200 μL S2 put all 10 tubes on ice mixed
- added 150 μL 7.5 M NH_4OAc
- incubated on ice for 5 min.
- cfg all 10 tubes for 5 min. @ room temp. (4°C)
- transferred 400 μL of supernate to the new 10 labelled tubes
- added 800 μL EtOH Mixed well
- incubated all 10 tubes for 30 min. @ -70°C .
- cfg & discard for 2 min. @ room temp.
- discarded supernate & washed pellet with 70% EtOH.
- added 50 μL TE to all 10 tubes w/ pellet

	tubes	
H_2O	1.0 μL x 10	= 10.0 μL
buffer	2.0 μL x 10	= 20.0 μL
ECO 47III	1.0 μL x 10	= 10.0 μL
TV		= 100.0 μL

- added 10.0 μL from TV to all other 9 tubes
- added 10.0 μL DNA to each 10 tubes
- incubated @ 37°C for 30 min.
- added 2 μL loading dye
- ran all 10 samples on a gel for 1 hour @ 190 V
- took a picture

picture on pg 35

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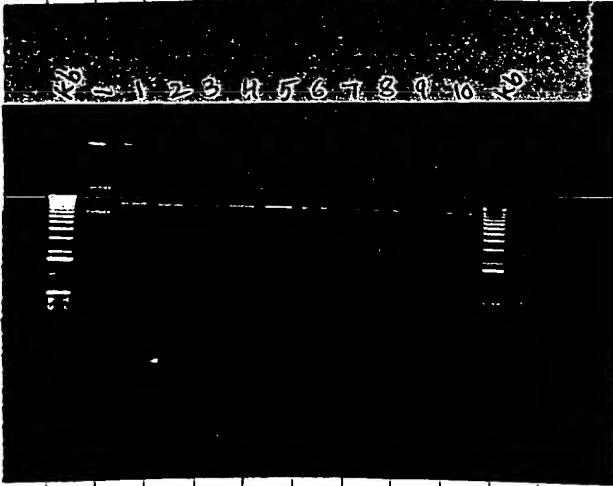
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parent & mutant should look like
ECO 47 III

	parent	mutant
8.9 kb	—	4.5 kb
0.3 kb	—	4.4 kb
		0.3 kb

3/16/95

1/12/95

may probably be too light to see

NOTE: In this ~~we~~ we could see parent & some mutant mutant is seen on # 5, 6, 7, 8

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